

Release Notes

caAMEL
Version 1.5
April 2007

National Cancer Institute Center for Bioinformatics

Contents

- 1.0 caAMEL Introduction and History
- 2.0 Release History
- 3.0 New Features and Updates
- 4.0 Bugs Fixed Since Last Release
- 5.0 Known Issues/Defects
- 6.0 Feature Requests
- 7.0 Bug Reports and Support
- 8.0 Documentation and Files
- 9.0 NCICB Web Pages

1.0 caAMEL Version 1.5 - Introduction and History

caAMEL is a web-based application that validates MAGE-ML documents, and more importantly, loads the data contained within valid MAGE-ML documents into a caArray repository. The caAMEL project, as well as the caArray project that it is targeted to enhance, is sponsored by the National Cancer Institute Center for Bioinformatics (NCICB).

The stated goals of NCICB's caArray application are "to make microarray data publicly available, and to develop and bring together open source tools to analyze these data." . Both of these objectives are predicated on caArray being widely adopted and on the population of caArray instances with a wealth of microarray experiment data. One key impediment in these areas has been the inability to import existing microarray design and experiment annotation into caArray. caAMEL is designed to fill this need by providing a mechanism to validate and load existing MAGE-ML documents into caArray.

caAMEL is a product of the NCI Center for Bioinformatics and its partners. Visit the caArray web site for more information:

-- <http://caarray.nci.nih.gov/>

2.0 Release History

caAMEL Version 1.5 (Major Release)
-- April 2007

caAMEL Version 1.5 RC1 (Major Release)
-- December 26 2006

caAMEL Version 1.0 (Major Release)
-- August 11 2006

3.0 New Features and Updates

- Tab and white space delimited data files associated with MAGE-ML documents are now automatically parsed and imported into caArray.

4.0 Defects Fixed Since Last Release

- See GForge Tracker

5.0 Known Issues/Defects

- See GForge Tracker

6.0 Feature Request

Feature requests may be submitted and reviewed at the
caAMEL GForge URL:

http://gforge.nci.nih.gov/tracker/?atid=557&group_id=118&func=browse

7.0 Bug Reports and Support

Send email to ncicb@pop.nci.nih.gov to request support or
report a bug. Existing bugs and resolution may be viewed
at the caAMEL GForge URL:

http://gforge.nci.nih.gov/tracker/?atid=554&group_id=118&func=browse

8.0 Documentation and Files

All documentation and files are bundled with the distribution.
Documentation can be found in the docs subdirectory of this
distribution.

9.0 NCICB Web Page

The NCI Center for Bioinformatics
-- <http://ncicb.nci.nih.gov/>

NCICB Application Support
-- <http://ncicbsupport.nci.nih.gov/sw/>

NCICB Download Center
-- <http://ncicb.nci.nih.gov/download/>

caArray
-- <http://caarray.nci.nih.gov/>